### In [1]:

```
import numpy as np
import pandas as pd
import seaborn as sns
from matplotlib import pyplot as plt
from sklearn.preprocessing import StandardScaler
from sklearn.model_selection import train_test_split
from sklearn.model_selection import cross_val_score
from sklearn import svm
from sklearn.model_selection import GridSearchCV
from keras.preprocessing.image import ImageDataGenerator
from PIL import Image
import PIL
import glob
import os
from keras.preprocessing.image import (
    ImageDataGenerator,
    load img,
    img to array,
    array_to_img
)
from keras.applications.resnet50 import ResNet50
from keras.applications.resnet50 import preprocess input,decode predictions
import skimage
from skimage import io
from skimage import transform
Using TensorFlow backend.
In [220]:
dataset = pd.read_csv("./Exercise1 - data.csv", header=None)
In [221]:
dataset.shape
Out[221]:
(569, 32)
In [222]:
dataset.iloc[:,1].value counts()
#the dataset is a little bit unbalanced
#Bening instances = 63%
#Metastatic instances = 37%
Out[222]:
В
     357
     212
Name: 1, dtype: int64
```

# Check if there are any null values

```
In [223]:
dataset.info()
# All Data are ok
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 569 entries, 0 to 568
Data columns (total 32 columns):
      569 non-null int64
1
      569 non-null object
2
      569 non-null float64
3
      569 non-null float64
4
      569 non-null float64
5
      569 non-null float64
      569 non-null float64
6
7
      569 non-null float64
      569 non-null float64
8
9
      569 non-null float64
10
      569 non-null float64
11
      569 non-null float64
      569 non-null float64
12
13
      569 non-null float64
14
      569 non-null float64
15
      569 non-null float64
16
      569 non-null float64
17
      569 non-null float64
      569 non-null float64
18
19
      569 non-null float64
      569 non-null float64
20
21
      569 non-null float64
22
      569 non-null float64
      569 non-null float64
23
24
      569 non-null float64
25
      569 non-null float64
      569 non-null float64
26
27
      569 non-null float64
28
      569 non-null float64
29
      569 non-null float64
      569 non-null float64
30
      569 non-null float64
dtypes: float64(30), int64(1), object(1)
memory usage: 142.3+ KB
In [224]:
```

```
# we can drop the ID column because I dont think we will need it
dataset = dataset.drop(dataset.columns[0], axis=1)
```

# Check if there is any type of correlation between features

### In [225]:

```
#considering that the feature are split into three big groups (means, standard errors an
d largest values)
#now we will consider just the group means and so the columns 3-12

mean_dataset = dataset.iloc[:,:11]
corr = mean_dataset.corr()
corr.style.background_gradient().set_precision(2)
```

### Out[225]:

	2	3	4	5	6	7	8	9	10	11
2	1	0.32	1	0.99	0.17	0.51	0.68	0.82	0.15	-0.31
3	0.32	1	0.33	0.32	-0.023	0.24	0.3	0.29	0.071	-0.076
4	1	0.33	1	0.99	0.21	0.56	0.72	0.85	0.18	-0.26
5	0.99	0.32	0.99	1	0.18	0.5	0.69	0.82	0.15	-0.28
6	0.17	-0.023	0.21	0.18	1	0.66	0.52	0.55	0.56	0.58
7	0.51	0.24	0.56	0.5	0.66	1	0.88	0.83	0.6	0.57
8	0.68	0.3	0.72	0.69	0.52	0.88	1	0.92	0.5	0.34
9	0.82	0.29	0.85	0.82	0.55	0.83	0.92	1	0.46	0.17
10	0.15	0.071	0.18	0.15	0.56	0.6	0.5	0.46	1	0.48
11	-0.31	-0.076	-0.26	-0.28	0.58	0.57	0.34	0.17	0.48	1

### In [226]:

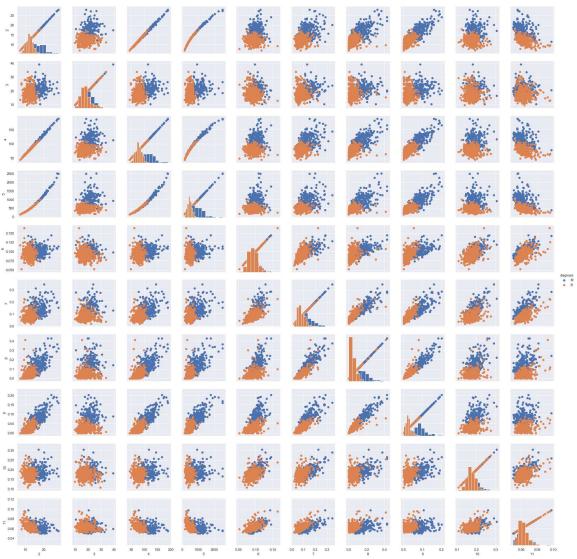
#between many variables there is a strong correlations, but to understand better #we will plot a PairGrid that will tell us if there is also a correlation with the diag nosis

### In [227]:

```
#rename the label column for semplicity
mean_dataset = mean_dataset.rename(columns={1:"diagnosis"})
```

### In [228]:

```
import matplotlib.pyplot as plt
import seaborn as sns;
g = sns.PairGrid(mean_dataset,hue="diagnosis")
g = g.map_diag(plt.hist)
g = g.map(plt.scatter)
g = g.add_legend()
```



### In [229]:

#How it is noticeable from this scatter plot, for the most variables (Mean values of ce ll radius, perimeter, area, compactness,

#concavity and concave points) if there are large values the diagnosis is metastatic.
#It means that there is a correlation between the most those mean variables and the diagnosis.

#Furthermore it looks like variables have gaussian distribution

# **Preprocessing Data**

### In [230]:

#Now we will prepare the data for the classification

### In [231]:

```
#First of all Transform the label that is a categorical variable into a numerical form

columns_label = pd.get_dummies(dataset[1])
dataset = dataset.drop(dataset.columns[0], axis=1)
dataset = pd.concat([dataset,columns_label], axis=1)
```

### In [262]:

```
input_data = dataset.iloc[:,:30]
#in label I will keep just the column (M) becasue it is a binary classification
label = dataset.iloc[:,31:]
```

### In [263]:

```
#Considering that the variable are Gaussian we can use the standardization method
# that standardize features by removing the mean and scaling to unit variance

scaler =StandardScaler()
input_standardize = scaler.fit_transform(input_data)
```

### In [264]:

#I was thinking about applying some dimensionality reduction methos as PCA but consider ed the size of the dataset it may be not #necessay

### **Prediction**

### In [259]:

#Considering the size of the dataset and that the the problem is a binary class one we decide to use SVM

### In [265]:

```
#transform Labels from pandas to numpy
label = label.values
label = label.ravel()

#The first thing to do is to split the dataset for training and test .
train_x, test_x, train_label, test_label = train_test_split(input_standardize, label, test_size=0.3)
```

### In [267]:

```
#Train a SVM on training set

clf = svm.SVC(probability=True)
clf.fit(train_x, train_label)

classifier_score = clf.score(test_x, test_label)
print ('\nThe classifier test accuracy score is {:.3f}\n'.format(classifier_score))
```

The classifier accuracy score is 0.982

### In [275]:

```
#Considering that it is not a time series another sophisticated way to calculate the ac
curacy
#is the cross validation, where basically the test set is not constant but it will chan
ge at each iteration

result = cross_val_score(svm.SVC(), input_standardize, label)

print (result)

#These three values represent the test accuracy, and are three becasue by default the n
umper of paritition created is 3.
#It means that there will be three iteration where 2 folds, namely 66% of data, will be
used for the training and the remaining
#one for the test.
```

[0.96315789 0.98421053 0.97354497]

In [277]:

```
max(result)
```

Out[277]:

0.9842105263157894

In [280]:

```
avg,uncertainty = result.sum()/3, [max(result)-result.sum()/3,result.sum()/3-min(result
)]
print("Average = {:2f}, uncertainty = [+{:2f},-{:2f}]".format(avg,uncertainty[0],uncertainty[1]))
```

Average = 0.973638, uncertainty = [+0.010573,-0.010480]

# Tuning hyper-parameters of a model

### In [282]:

```
#Tuning the hyper-parameters of a model is a fundamental partthat may help to improve t
he performance
#One easy way to do that in python is using GridSearch
svm.SVC().get params()
#The most important parameter that can be tuned are:
#- C that represent the cost of misclassification
#- s the parameter of a Gaussian Kernel
#- kernel says the type of kernel used to classify the points, the rbf uses normal curv
es around the data points
Out[282]:
{'C': 1.0,
 'cache size': 200,
 'class_weight': None,
 'coef0': 0.0,
 'decision function shape': 'ovr',
 'degree': 3,
 'gamma': 'auto',
 'kernel': 'rbf',
 'max iter': -1,
 'probability': False,
 'random_state': None,
 'shrinking': True,
 'tol': 0.001,
 'verbose': False}
In [292]:
kernel_types = [ 'linear' , 'poly' ,
                                        'rbf' , 'sigmoid' ]
C = [0.00001, 0.0001, 0.01, 0.1, 1, 10]
gamma = [0.001, 0.01, 0.1, 1]
param_grid = {'C': C, 'gamma' : gamma, 'kernel':kernel_types}
grid_search = GridSearchCV(svm.SVC(), param_grid)
grid_search.fit(train_x, train_label)
print("Best parmas are : " , grid_search.best_params_)
print("Best score is : ", grid_search.best_score_)
Best parmas are : {'gamma': 0.001, 'kernel': 'linear', 'C': 0.1}
Best score is: 0.9748743718592965
In [294]:
#Train a SVM on training set with best parmas
clf = svm.SVC(probability=True, C=0.1, gamma=0.001, kernel='linear')
clf.fit(train_x, train_label)
classifier score = clf.score(test x, test label)
print ('\nThe classifier test accuracy score is {:.3f}\n'.format(classifier_score))
```

The classifier test accuracy score is 0.988

# Classification using raw images

### In [295]:

```
#The easiest way to do this is using some pre-trained model as ResNet but first of all
I will try to build
# a deep network from scratch just to demonstrate that I know what I am talking about
```

### In [54]:

```
#convert from GIF to jpg
files = glob.glob("./Raw_images/Training/Training_Malignant/*.gif")
# print(files)
for imageFile in files:
    filepath,filename = os.path.split(imageFile)
    filterame,exts = os.path.splitext(filename)
    print ("Processing: " + imageFile,filterame)
    im = Image.open(imageFile)
    if im.mode in ['1', 'P']:
        im = im.convert('RGB')
    #increase number of image (augmentation)
    #rotate
    img rotated 90 = im.rotate( angle=90)
    img rotated 180 = im.rotate( angle=180)
    im.save( './Raw images/Training/Train Benign'+filterame+'.jpg','JPEG')
    img_rotated_90.save( './Raw_images/Training_good/Train_Benign/'+filterame+'_rotated
_90.jpg','JPEG')
    img_rotated_180.save( './Raw_images/Training_good/Train_Malign/'+filterame+'_rotate
d 180.jpg','JPEG')
Processing: ./Raw images/Training/Training Malignant\91 6838.gif 91 6838
Processing: ./Raw_images/Training/Training_Malignant\91_9555.gif 91_9555
Processing: ./Raw images/Training/Training Malignant\92 6125.gif 92 6125
Processing: ./Raw_images/Training/Training_Malignant\92_6424.gif 92_6424
```

### In [57]:

```
from keras.models import Sequential
from keras.layers import Conv2D
from keras.layers import MaxPooling2D
from keras.layers import Flatten
from keras.layers import Dense
```

Processing: ./Raw\_images/Training/Training\_Malignant\92\_6682.gif 92\_6682 Processing: ./Raw\_images/Training/Training\_Malignant\92\_6954.gif 92\_6954 Processing: ./Raw images/Training/Training Malignant\92\_7241.gif 92\_7241

```
In [58]:
```

```
train_generator = ImageDataGenerator(rescale = 1./255,
shear_range = 0.2,
zoom_range = 0.2,
horizontal_flip = True)
test_generator = ImageDataGenerator(rescale = 1./255)
training_set = train_generator.flow_from_directory('./Raw_images/Training_good',
target_size = (64, 64),
class_mode = 'binary')
test_set = test_generator.flow_from_directory('./Raw_images/Test_good',
target_size = (64, 64),
class_mode = 'binary')
```

Found 37 images belonging to 2 classes. Found 5 images belonging to 2 classes.

### In [64]:

```
# Initialising the CNN
classifier = Sequential()
# Step 1 - Convolution
classifier.add(Conv2D(64, (3, 3), input shape = (64, 64, 3), activation = 'relu'))
# Step 2 - Pooling
classifier.add(MaxPooling2D(pool size = (2, 2)))
# Adding a second convolutional layer
classifier.add(Conv2D(32, (2, 2), activation = 'relu'))
classifier.add(MaxPooling2D(pool size = (2, 2)))
# Step 3 - Flattening
classifier.add(Flatten())
# Step 4 - Full connection
classifier.add(Dense(units = 128, activation = 'relu'))
classifier.add(Dense(units = 1, activation = 'sigmoid'))
# Compiling the CNN
classifier.compile(optimizer = 'adam', loss = 'binary crossentropy', metrics = ['accura
cy'])
classifier.fit_generator(training_set,
steps_per_epoch = 37,
epochs = 2,
validation data = test set,
validation_steps = 5)
```

# **Using ResNet**

<keras.callbacks.History at 0x18c8ae99898>

```
In [ ]:
model=ResNet50(include_top=True, weights='imagenet')
Downloading data from https://github.com/fchollet/deep-learning-models/rel
eases/download/v0.2/resnet50_weights_tf_dim_ordering_tf_kernels.h5
 36806656/102853048 [=======>.....] - ETA: 8:16
In [ ]:
def hack_resnet(input_size):
    base_model = ResNet50(include_top=False, weights='imagenet', input_shape=input_size
)
    x = base model.output
    x = Flatten()(x)
    x = Dense(num_classes, activation='sigmoid', name='fc1000')(x)
    # this is the model we will train
    newmodel = Model(inputs=base model.input, outputs=x)
    return newmodel
model = hack_resnet(training_set.image_shape)
In [ ]:
```

### **Final Consideration**

In this challenge I tried to predict if a patient had a cancer or not. In first place I tried SVM due to the size of the dataset, namely 569 rows and after plotting some statistics I built and SVM model that performed very well, moreover I improved a little bit the performance tuning the parameters and reaching an 98% on the test set. After I tried to do the same using raw images. I used a CNN network. Due to the small number of images I applied some augmentation methods. I reached an accuracy of 80% that consider the small number of validation images (5) meanas 4 out of 5 classified correctly. After I wanted to try transfer learning but due to internet problem i was not able to download the pre-trained model.

mode.compile(loss='binary\_crossentropy',optimizer = 'adam', metrics=['accuracy'])